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INFORMATION DISCLOSURE STATEMENT BY APPLICANT (use as many sheets as necessary)		Application Number	10/009,876
		Filing Date	December 11, 2001
		First Named Inventor	R.G.F. Visser
		Group Art Unit	TBA
		Examiner Name	TBA
Sheet	2	of	2
		Attorney Docket Number	92750/65

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Examiner Initials	Cite No. ¹	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published	T ²
AK	11	CHEN et al., Improved Adsorption to Starch of a Beta-Galactosidase Fusion Protein Containing the Starch-Binding Domain from Aspergillus Glucoamylase. Biotechnology Progress, 7:225-29, 1991	
I	12	DALMIA et al., Domain E of Bacillus macerans Cyclodextrin Glucanotransferase: An Independent Starch-Binding Domain. Biotechnology and Bioengineering, 47:575-84, 1995	
	13	KUSNADI et al., Functional Starch-Binding Domain of Aspergillus Glucoamylase I in Escherichia Coli. Gene, 127(2):193-97, 1993	
	14	LAWSON et al., Nucleotide Sequence and X-ray Structure of Cyclodextrin Glycosyltransferase from Bacillus circulans Strain 251 in a Maltose-dependent Crystal Form. J. Mol. Biol., 236:590-600, 1994	
	15	OHDAN et al., Introduction of Raw Starch-Binding Domains into Bacillus subtilis Alpha-Amylase by Fusion with the Starch-Binding Domain of Bacillus Cyclomaltodextrin Glucanotransferase. Appl. Environ Microbiol., 66(7):3058-64, 2000	
	16	ONG et al., The Cellulose-Binding Domains of Cellulases: Tools for Biotechnology. Trends in Biotechnology, 7:239-43, 1989	
	17	PENNINGA et al., The Raw Starch Binding Domain of Cyclodextrin Glycosyltransferase from Bacillus circulans Strain 251. J. Biol. Chem, 271(51):32777-32784, 1996	
	18	SORIMACHI et al., Solution Structure of the Granular Starch Binding Domain of Glucoamylase from Aspergillus niger by Nuclear Magnetic Resonance Spectroscopy. J. Mol. Biol., 259(5):970-87, 1996	
	19	SORIMACHI et al., Solution Structure of the Granular Starch Binding Domain of Aspergillus niger Glucoamylase Bound to Beta-Cyclodextrin. Structure, 5(5):647-61, 1997	
AK	20	SVENSSON et al., Sequence homology between putative raw-starch binding domains from different starch-degrading enzymes. Biochem. J., 264:309-11, 1989	

Examiner Signature	Russell K Allen	Date Considered	5/5/05
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